CLAIMS

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1. A purified polynucleotide comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO:2, or the complement of said polynucleotide.

The polynucleotide of Claim 1 comprising the nucleic acid sequence of 2.

pdynucleatide omprising the complement of the 3. polynucleotide of Claim 2 or a portion thereof.

A pharmaceutical composition comprising the antisense molecule of Claim 3 and a pharmaceutically acceptable excipient.

composition comprising an oligomer of the polynucleotide 5. of Claim 2.

A diagnostic test for a condition associated with altered VR-L 6. expression comprising the steps of:

providing a biological sample; a)

- combining the biological sample and the diagnostic composition b) of Claim 5
- c) allowing hybridisation to occur between the biological sample and the diagnostic composition under suitable conditions;
- d) measuring the amount of hybridisation to obtain a sample value; and
 - comparing the sample value with standard values to determine e) whether vr-l expression is altered.
- 7. An expression vector comprising the polynucleotide of Claim 1.
- 25 8. A host cell transformed with the expression vector of Claim 7.
 - 9. A method of producing a polypeptide, said method comprising the steps of:
 - a) culturing the host cell of Claim 8 under conditions suitable for the expression of the polypeptide; and

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- b) recovering the polypeptide from the host cell culture.
- 10. A purified polypeptide (VR-L) comprising the amino acid sequence of SEQ ID NO:2.
- 11. A diagnostic composition comprising the polypeptide of Claim 10 or a portion thereof.
 - 12. A pharmaceutical composition comprising the polypeptide of Claim 10 and a pharmaceutically acceptable excipient.
 - 13. An antibody specific for the purified polypeptide of Claim 9, or for a portion of that polypeptide.
 - 14. A diagnostic composition comprising the antibody of Claim 13.
 - 15. A diagnostic test for a condition associated with altered VR-L expression comprising the steps of:
 - a) providing a biological sample;
 - b) combining the biological sample and the antibody of Claim 13 under conditions suitable for complex formation;
 - c) measuring the amount of complex formation between VR-L and the antibody to obtain a sample amount; and
 - d) comparing the amount of complex formation in the sample with standard amounts of complex formation, wherein a variation between sample amount and standard amounts of complex formation establishes the presence of the condition.
 - 16. A method of screening a plurality of compounds for specific binding affinity with the polypeptide of Claim 10 or any portion thereof comprising the steps of:
 - a) providing a plurality of compounds;
 - b) combining VR-L with each of a plurality of compounds for a time sufficient to allow binding under suitable conditions; and
 - c) detecting binding of VR-L to each of the plurality of compounds, thereby identifying the compounds which specifically bind VR-L.

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17. A pharmaceutical composition comprising a compound of Claim 16 and capharmaceutically acceptable excipient.